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	the state of the s	Fig. 1 (2011) The first from the fir		
The state of the s	*Vergleich Endometriose 4	VerglejchEndometriose	Vergleich sekrit Versus prolitikase	
	versus Normal (sekr. Phase)	versusiNormali (prolinghase)		
Control (in the precise of the preci	down (0 up - 16 down)	down (4 up -12 down)	up (18 up - 1 down)	
x02701, included arowth factor binding protein-2	down (1-15)	nc (13-13)	up (17-2)	
U40271, Human transmembrane receptor precursor	down (0-14)	nc (6-2)	up (9-1)	
(PTK7)				
M21574, platelet-derived growth factor receptor alpha	down (0-13)	nc (8-10)	up (17-0)	1/1-
(PDGFRA)			(17-0)	
1 22548 collagen type XVIII alpha 1 (COL18A1)	down (0-13)	down (0-8)	(n-11) dn	
	down (1-13)	down (4-13)	up (22-2)	
M80482, subtilisin-like protein (PACE4)			(17.4)	
226653, laminin M chain (merosin)	down (1-13)	nc (9-10)	() dn	
The state of the s	down (0-12)	nc (0-0)	up (25-0)	
M36860, U77840, Elastiii			up (11-0)	
X05610, type IV collagen alpha -2 chain	down (0-12)	nc (3-3)		 -X
X67325, p27 interferon alpha-inducible gene	down (1-12)	nc (9-10)	nb (10-7)	

Abbildung 1

[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name

Vergleich Endometriose versus Normal (sekr. Phase) = Comparison of Endometriosis versus Normal (Secr. Phase)

Vergleich Endometriose versus Normal (prol. Phase) = Comparison of Endometriosis versus Normal (prol. Phase)

Vergleich sekr. versus prol. Phase (Endometrium) = Comparison of Secr. versus Prol. Phase (Endometrium)

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Datenbank-Nr., Name	Vergleich Endometriose	Vergleich Endometriose	Vergleich sekr. versus prol. Phase
	versus Normal (sekr. Phase)	versus Normal (prol. Phase)	(Endometrium)
D42073, reticulocalbin	down (0-11)	nc (8-5)	up (11-2)
U07919, aldehyde dehydrogenase 6	down (1-11)	nc (13-9)	up (22-0)
U81607, gravin	down (1-11)	nc (8-7)	up (18-1)
M30269, nidogen	down (0-10)	nc (8-14)	up (15-3)
D42108, phospholipase C Epsilon	down (1-10)	nc (12-14)	up (25-0)

Abbildung 1

1

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[Key to Table:]

Datenbank-Nr., Name = Data Bank No., Name

Vergleich Endometriose versus Normal (sekr. Phase) = Comparison

of Endometriosis versus Normal (prol. Phase) = Comparison

of Endometriose versus Normal (prol. Phase) = Comparison

of Endometriosis versus Normal (prol. Phase)

Vergleich sekr. versus prol. Phase (Endometrium) = Comparison

of Secr. versus Prol. Phase (Endometrium)

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Abbildung 2		
Seq.IDNO	Name	Proteinsequenz
	Fibronektin	VPSTGASKSK RQAQQMVQPQ SPVAVSQSKP GCIDNGKHIV
•		GSRGFNCESK PEAEEICFUN 11GN118000
		GDIWKRFREI GOTHELSTEIN EIGDTWSKKD NRGNLLQCIC TGNGRGEWKC
		OPPPYGHCVT DSGVVYSVGM QMLKTQGNKQ MLCTCLGNGV
		VLPFTYNGRT FYSCTTEGRQ DGHLWCSTTS NYEQDQKYSF CTDHTVLVQT
		YTDCTSEGRR DNMKWCGTTQ NYDADQKFGF CPMAAHEEIC TTNEGVMYRI
	-	MMRCTCVG NGRGEWTCIA YSQLRDQCIV DDITYNVNDT FHKRHEEGHM LNCTCFGQGR
		QIGDSWEKYV HGVRYQCYCY GRGIGEWHCQ PLQTYPSSSG PVEVFITETP
		YILRWRPKNS VGRWKEATIP GHLNSYTIKG LKPGVVYEGQ LISIQQYGHU
		VTGETTPFSP LVATSESVTE ITASSFVVSW VSASDTVSGF
and the second seco		MENNION LEGRETION YOISEDGEQS LILSTSOTTA PDAPPDFTVD QVDDTSIVVR
		CSCTELAL OF TANSVILSDL QPGVQYNITI YAVEENQEST PVVIQQETTG
		VICTORIANGED PSAVIGYRVD VIPVNLPGEH GORLPISRNT FAEVTGLSPG
		DLQFVEVID VNVII:
		HGRESKPLI ACCITIONING TO THE CONTROL OF THE CONTROL
		PLKNLUPAS EIIVSLYIII SKATTOOLENDE ERDAPIVNKV VTPLSPPTNL
		POLITICARITY TPINGOOGNS LEEVVHADOS SCTFDNLSPG LEYNVSVYTV
		TOTAL BETWIED DIMRVIMAPP PSIDLINFLY RYSPVKNEED VAELSISPSD
		VAROHRSTPL RGROKTGLDS PTGIDFSDIT ANSFTVHWIA PRATITGYRI
		GIEIVVSVSS VIEWARD VITTIANLIP GIEWVSIVA LNGREESPLL IGQQSTVSDV PRDLEVVAAT PTSLLISWDA

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[Key to Table:]

Proteinsequenz = Protein Sequence Fibronektin = fibronectin

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Seq.IDNO	Name	Proteinsequenz
		ļ
,		SQPLVQTAVT NIDRPKGLAF TDVDVDSIKI AWESPQGQVS RYRVTYSSPE
		GLRPGSEYTV SVVALHDDME SQPLIGTQST AIPAPTDLKF TQVTPTSLSA
		RVRVTPKE KTGPMKEINL APDSSSVVVS GLMVATKYEV SVYALKDTLT SRPAQGVVTT
		TDATETTI TISWRTKTET ITGFQVDAVP ANGQTPIQRT IKPDVRSYTI TGLQPGTDYK
		TAIDAPSNLR FLATTPNSLL VSWQPPRARI TGYIIKYEKP GSPPREVVPR
		LEPGTEYT IYVIALKNNQ KSEPLIGRKK TDELPQLVTL PHPNLHGPEI LDVPSTVQKT
		GOOPSVGQQM IFEEHGFRRT TPPTTATPIR HRPRPYPPNV GEEIQIGHIP
		CHRRCHSERW CHDNGVNYKI GEKWDRQGEN GQMMSCTCLG NGKGEFKCDP HEATCYDDGK TYHVGEQWQK
		CFGGQRGWRC
		REDSRE
2	Insulin-like	VIEWCEECE
1	growth factor	AGGARMPCAE LVREPGCGCC SVCARLEGEA CGVYTPRCGQ GLRCYPHPGS ELPHAALVWG
	binding protein-2	EYGASPEQVA DNGDDHSEGG LVENHVDSTM NMLGGGGSAG RKPLKSGMKE LAVFREKVTE QHKQMGKGGK
· · · · · · · · · · · · · · · · · · ·		HHLGLEEPKK LRPPPARTPC QQELDQVLER ISTMRLPDER GPLEHLYSLH IPNCDKHGLY NLKQCKMSLN

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[Key to Table:]
Proteinsequenz = Protein Sequence

Seq.IDNO	Name	Proteinsequenz
	Transmembrane	MGAARGSPAR PRRLPLLSVL LLPLLGGTQT AIVFIKQPSS QDALQGRRAL LRCEVEAPGP VHVYWLLDGA
n	receptor PTK7	
	1 1 1 1 1 1 1 1 1	
		SSQNFTLSIA DESFARVVLA PQDVVVARYE EAMFHCQFSA QPPPSLQWLF EDETPITNRS RPPHLRRATV
www.to	-	FANGSLLLTQ VRPRNAGIYR CIGQGQRGPP IILEATLHLA EIEDMPLFEP RVFTAGSEER VTCLPPKGLP
		EPSVWWEHAG VRLPTHGRVY OKGHELVLAN IAESDAGVYT CHAANLAGOR RODVNITVAT VPSWLKKPOD
		SQLEEGKPGY LDCLTQATPK PTVVWYRNQM LISEDSRFEV FKNGTLRINS VEVYDGTWYR CMSSTPAGSI
		EAQAVLQVLE KLKFTPPPQP QQCMGFDKEA TVPCSATGRE KPTIKWERAD GSSLPEWVTD NAGTLHFARV
		TRDDAGNYTC IASNGPQGQI RAHVQLTVAV FITFKVEPER TTVYQGHTAL LQCEAQGDPK PLIQWKGKDR
		ILDPTKLGPR MHIFQNGSLV IHDVAPEDSG RYTCIAGNSC NIKHTEAPLY VVDKPVPEES EGPGSPPPYK
		MIOTIGLSVG AAVAYIIAVL GLMFYCKKRC KAKRLQKQPE GEEPEMECLN GGPLQNGQPS AEIQEEVALT
		SLGSGPAATN KRHSTSDKMH FPRSSLQPIT TLGKSEFGEV FLAKAQGLEE GVAETLVLVK SLQSKDEQQQ
		LDFRRELEMF GKLNHANVVR LLGLCREAEP HYMVLEYVDL EDLKQFLRIS KSKDEKLKSQ PLSTKQKVAL
		CTQVALGMEH LSNNRFVHKD LAARNCLVSA QRQVKVSALG LSKDVYNSEY YHFRQAWVAL RWMSPEAILE
		GDFSTKSDVW ASGVLMWEVF THGEMPHGGQ ADDEVLADLQ AGKARLPQPE GCPSKLYRLM ORCWALSPKD
-		RPSFSEIASA LGDSTVDSKP
7	Platelet-derived	MGTSHPAFLV LGCLLTGLSL ILCQLSLPSI LPNENEKVVQ LNSSFSLRCF GESEVSWQYP MSEEESSDVE
 	growth factor	VPDPDVAFVP
	receptor alpha	EDDDSAIIPC RTTDPETPVT LHNSEGVVPA SYDSRQGFNG TFTVGPYICE ATVKGKKFQT IPFNVYALKA
	4 1 1 2 3 3 4 3 3 3 4	

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[Key to Table:]
Proteinsequenz = Protein Sequence

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Seq.IDNO	Name	Proteinsequenz
		TVKDSGDYEC AARQATREVK EMKKVTISVH EKGFIEIKPT FSQLEAVNLH EVKHFVVEVR AYPPFKISWL
		KANLTLIENL TEITTDVEKI QEIRYRSKLK LIRAKEEDSG HYTIVAQNED AVKSYTFELL TQVPSSILDL
		UNDHHGSTGG OTVRCTAEGT PLPDIEWMIC KDIKKCNNET SWIILANNVS NIITEIHSRD RSTVEGRVTF
	••	PRSSEKOA
		PKKELDIFGL NPADESTRSY VILSFENNGD YMDMKQADTT QYVPMLERKE VSKYSDIQRS LYDRPASYKK
		KEMINSEVKN LLSDDNSEGL TLLDLLSFTY QVARGMEFLA SKNCVHRDLA ARNVLLAQGK IVKICDFGLA
		SKGSTFLPVK
· .		VDSDNAYIG VTYKNE
		SSOTSEESAI ETGSSSSTFI KREDETIEDI DMMDDIGIDS SDLVEDSFL
	type	GEVGADGIPG FPGLPGREGI AGPQGPKGDR GSRGEKGDPG KDGLGQPGLP GPRGPPGPVV
<u>.</u>	nha 1	SVPGPEGRRG FAGFPGPAGP KGNLGSKGEL GSPGPKGEKG EPGSIFSPDG GALGPAQKGA
		GLYGRPGYKG EIGFPGRPGR PGMNGLKGEK GEPGDASLGF GMRGMPGPPG PPGPPGPPGT PVYDSNVFAE
		SSRPGPPGLP GNQGPPGPKG PKGEVGPPGP PGQFPFDFLQ KEAEMKGEKG DRGDAGQKGE RGEPGGGGFF
		SSLPGAPGA
		PPGPPGTMGA SSGQVRLWAT RQAMLGQVHE VPEGWLIFVA EQEELYVRVQ
		RTPLPRGTON EVAALQPPVV QLHDSNPYPR REHPHPTARP WRADDILASP PGLPEPQPYP GGPHHSSYVH

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		IVRRADRAAV PIVNLKDELL FPSWEALFSG SEGPLKPGAR IFSFDGKDVL RHPTWPQKSV WHGSDPNGRR LTESYCETWR TEAPSATGQA SSLLGGRLLG QSAASCHHAY IVLCIENSFM TASK
	Subtiliain-like	MPPRAPPAPG PRPPPRAAAA TDTAAGAGGA GGAGGAGGPG FRPLAPRPWR WLLLLALPAA CSAPPPRPVY
0	protein (PACE4	
,		QEVKRRVKRQ VRSDPQALYF NDPIWSNMWY LHCGDKNSRC RSEMNVQAAW KRGYTGKNVV VTILDDGIER
		NHPDLAPNYD SYASYDVNGN DYDPSPRYDA SNENKHGTRC AGEVAASANN SYCIVGIAYN AKIGGIRMLD
		GDVTDVVEAK SLGIRPNYID IYSASWGPDD DGKTVDGPGR LAKQAFEYGI KKGRQGLGSI FVWASGNGGR
		EGDYCSCDGY TNSIYTISVS SATENGYKPW YLEECASTLA TTYSSGAFYE RKIVTTDLRQ RCTDGHTGTS
		VSAPMVAGII ALALEANSOL TWRDVQHLLV KTSRPAHLKA SDWKVNGAGH KVSHFYGFGL VDAEALVVEA
		KKWTAVPSQH MCVAASDKRP RSIPLVQVLR TTALTSACAE HSDQRVVYLE HVVVRTSISH PRRGDLQIYL
		VSPSGTKSQL LAKRLLDLSN EGFTNWEFMT VHCWGEKAEG QWTLEIQDLP SQVRNPEKQG KLKEWSLILY
		GTAEHPYHTF SAHQSRSRML ELSAPELEPP KAALSPSQVE VPEDEEDYTA QSTPGSANIL QTSVCHPECG
		DKGCDGPNAD QCLNCVHFSL GSVKTSRKCV SVCPLGYFGD TAARRCRRCH KGCETCSSRA ATQCLSCRRG
		FYHHQEMNTC VTLCPAGFYA DESQKNCLKC HPSCKKCVDE PEKCTVCKEG FSLARGSCIP DCEPGTYFDS
		ELIRCGECHH TCGTCVGPGR EECIHCAKNF HFHDWKCVPA CGEGFYPEEM PGLPHKVCRR CDENCLSCAG
		SSRNCSRCKT GFTQLGTSCI TNHTCSNADE TFCEMVKSNR LCERKLFIQF CCRTCLLAG
1	Caminin M chain	Chain MPGAAGVLLL LLLSGGLGGV QAQRPQQRQ SQAHQQRGLF PAVLNLASNA LITTNATCGE KGPEMYCKLV
	:	EHVPGQPVRN PQCRICNQNS SNPNQRHPIT NAIDGKNTWW QSPSIKNGIE YHYVTITLDL QQVFQIAYVI
		VKAANSPRPG NWILERSLDD VEYKPWQYHA VTDTECLTLY NIYPRTGPPS YAKDDEVICT SFYSKIHPLE
		NGEIHISLIN GRPSADDPSP ELLEFTSARY IRLRFORIRT LNADLMMFAH KDPREIDPIV TRRYYYSVKD
		ISVGGMCICY GHARACPLDP ATNKSRCECE HNTCGDSCDQ CCPGFHQKPW RAGTFLTKTE CEACNCHGKA

[Key to Table:]
Proteinsequenz = Protein Sequence

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Seq.IDNO	Name	Proteinsequenz
,		EECYYDENVA RRNLSLNIRG KYIGGGVCIN CTQNTAGINC ETCTDGFFRP KGVSPNYPRP COPCHCDPIG
		SLNEVCVKDE KHARRGLAPG SCHCKTGFGG VSCDRCARGY TGYPDCKACN CSGLGSKNED PCFGPCICKE
		NVEGGDCSRC KSGFFNLQED NWKGCDECFC SGVSNRCQSS YWTYGKIQDM SGWYLTDLPG RIRVAPQQDD
		LDSPQQISIS NAEARQALPH SYYWSAPAPY LGNKLPAVGG QLTFTISYDL EEEEEDTERV LQLMIILEGN
		DLSISTAQDE VYLHPSEEHT NVLLLKEESF TIHGTHFPVR RKEFMTVLAN LKRVLLQITY SFGMDAIFRL
		SSUNLESAUS YPTDGSIAAA VEVCQCPPGY TGSSCESCWP RHRRVNGTIF GGICEPCQCF GHAESCDDVT
		GECLNCKDHT GGPYCDKCLP GFYGEPTKGT SEDCQPCACP LNIPSNNFSP TCHLDRSLGL ICDGCPVGYT
		GPRCERCAEG YFGQPSVPGG SCQPCQCNDN LDFSIPGSCD SLSGSCLICK PGTTGRYCEL CADGYFGDAV
		DAKNCOPCRC NAGGSFSEVC HSQTGQCECR ANVQGQRCDK CKAGTFGLQS ARGCVPCNCN SFGSKSFDCE
		GOCWCOPG
		HSITTGCKAC NCSTVGSLDF QCNVNTGQCN CHPKFSGAKC TECSRGHWNY PRCNLCDCFL PGTDATTCDS
		ETKKCSCSDO TGQCTCKVNV EGIHCDRCRP GKFGLDAKNP LGCSSCYCFG TTTQCSEAKG LIRTWVTLKA
		LILPLVDE
		EETGESTY
		LYDIHYIL
		SOPGGRTP
		PVTGFCTCRP GATGRKCDGC KHWHAREGWE CVFCGDECTG LLLGDLARLE OMVMSINLTG PLPAPYKMLY
		GLENMTQELK HLLSPQRAPE RLIQLAEGNL NTLVTEMNEL LTRATKVTAD GEQTGQDAER TNTRAKSLGE
		FIKELARDAE AVNEKAIKLN ETLGTRDEAF ERNLEGLQKE IDQMIKELRR KNLETQKEIA EDELVAAEAL
		LKKVKKLFGE SRGENEEMEK DLREKLADYK NKVDDAWDLL REATDKIREA NRLFAVNQKN MTALEKKKEA
		VESGKRQIEN TLKEGNDILD EANRLADEIN SIIDYVEDIQ TKLPPMSEEL NDKIDDLSQE IKDRKLAEKV

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Seq.IDNO	Name	Proteinsequenz
		SOAESHAAOL NDSSAVLDGI LDEAKNISFN ATAAFKAYSN IKDYIDEAEK VAKEAKDLAH EATKLATGPR
		GLLKEDAKGC LQKSFRILNE AKKLANDVKE NEDHLNGLKT RIENADARNG DLLRTLNDTL GKLSAIPNDT
		AAKLQAVKDK ARQANDTAKD VLAQITELHQ NLDGLKKNYN KLADSVAKTN AVVKDPSKNK IIADADATVK
		NLEQEADRLI DKLKPIKELE DNLKKNISEI KELINQARKQ ANSIKVSVSS GGDCIRTYKP EIKKGSYNNI
		VVNVKTAVAD NLLFYLGSAK FIDFLAIEMR KGKVSFLWDV GSGVGRVEYP DLTIDDSYWY RIVASRTGRN
		GTISVRALDG PKASIVPSTH HSTSPPGYTI LDVDANAMLF VGGLTGKLKK ADAVRVITFT GCMGETYFDN
		KPIGLWNFRE KEGDCKGCTV SPQVEDSEGT ATRDLRDFMS VELTDGHIKV SYDLGSGMAS VVSNQNHNDG
		KWKSFTLSRI QKQANISIVD IDTNQEENIA TSSSGNNFGL DLKADDKIYF GGLPTLRNLS MKARPEVNLK
		KYSGCLKDIE ISRTPYNILS SPDYVGVTKG CSLENVYTVS FPKPGFVELS PVPIDVGTEI NLSFSTKNES
		GIILLGSGGT PAPPRRKRRQ TGQAYYVILL NRGRLEVHLS TGARTMRKIV IRPEPNLFHD GREHSVHVER
***************************************		TRGIFTVQVD ENRRYMQNLT VEQPIEVKKL FVGGAPPEFQ PSPLRNIPPF EGCIWNLVIN SVPMDFARPV
		SFKNADIGRC AHOKLREDED GAAPAEIVIQ PEPVPTPAFP TPTPVLTHGP CAAESEPALL IGSKQFGLSR
		NSHIAIAFDD TKVKNRLTIE LEVRTEAESG LLFYMAAINH ADFATVQLRN GLPYFSYDLG SGDTHTMIPT
		KINDGQWHKI KIMRSKQEGI LYVDGASNRT ISPKKADILD VVGMLYVGGL PINYTTRRIG PVTYSIDGCV
		RNLHMAEAPA DLEQPTSSFH VGTCFANAQR GTYFDGTGFA KAVGGFKVGL DLLVEFEFAT TTTTGVLLGI
		SSOKMDGMGI EMIDEKLMFH VDNGAGRFTA VYDAGVPGHL CDGQWHKVTA NKIKHRIELT VDGNQVEAQS
		PNPASTSADT NDPVFVGGFP DDLKQFGLTT SIPFRGCIRS LKLTKGTASH WRLILPRPWN
8	Elastin	MAGLTAAAPR PGVLLLLLSI LHPSRPGGVP GAIPGGVPGG VFYPGAGLGA LGGGALGPGG KPLKPVPGGL
		AGAGLGAGLG AFPAVTFPGA LVPGGVADAA AAYKAAKAGA GLGGVPGVGG LGVSAGAVVP QPGAGVKPGK
		VPGVGLPGVY PGGVLPGARF PGVGVLPGVP TGAGVKPKAP GVGGAFAGIP GVGPFGGPQP GVPLGYPIKA
		PKLPGGYGLP YTTGKLPYGY GPGGVAGAAG KAGYPTGTGV GPQAAAAAAA KAAAKFGAGA AGVLPGVGGA

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Seq.IDNO	Name	Proteinsequenz	dnenz					
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		GVPGVPGAIP	GIGGIAGVGT PAA	AAAAAA	AKAAKYGAAA	PAAAAAAAA AKAAKYGAAA GLVPGGPGFG PGVVGVPGAG VPGVGVPGAG	PGVVGVPGAG	VPGVGVPGAG
		IPVVPGAGIP	GAAVPGVVSP	EAAKAAAKA 1	AKYGARPGVG	VGGIPTYGVG AGGFPGFGVG	AGGFPGFGVG	VGGIPGVAGV
		PSVGGVPGVG	GVPGVGISPE	AAAAKAA I	CYGVGTPAAA	AQAAAAKAA KYGVGTPAAA AAKAAAKAAQ	FALLNLAGLV	PGVGVAPGVG
-		VAPGVGVAPG	VGLAPGVGVÄ	PGVGVAPGVG V	/APGIGPGGV	VAPGIGPGGV AAAAKSAAKV AAKAQLRAAA	aakaqlraaa	GLGAGIPGLG
		VGVGVPGLGV	GAGVPGLGVG	AGVPGFGAVP (SALAAAKAAK	GALAAAKAAK YGAAVPGVLG GLGALGGVGI	GLGALGGVGI	PGGVVGAGPA
		AAAAAAKAAA	KAAQFGLVGA AGLGGLGVGG LGVPGVGGLG GIPPAAAAKA AKYGAAGLGG VLGGAGQFPL	3GLGVGG 1	GVPGVGGLG	GIPPAAAAKA	AKYGAAGLGG	VLGGAGQFPL
		GGVAARPGFG	LSPIFPGGAC	LGKACGRKRK				
6	Alpha-2 type IV	MGRDQRAVAG	PALRRWLLLG TVTVGFLAQS VLAGVKKFDV	VGFLAQS		PCGGRDCSGG	CQCYPEKGGR	GQPGPVGPQG
	collagen	YNGPPGLQGF	PGLQGRKGDK	GERGAPGVTG	PKGDVGARGV	SGFPGADGIP	GHPGQGGPRG	RPGYDGCNGT
		QGDSGPQGPP	GSEGFTGPPG PQGI	PQGPKGQKGE	PYALPKEERD	RYRGEPGEPG	LVGFQGPPGR	PGHVGQMGPV
		GAPGRPGPPG	PPGPKGQQGN RGL	RGLGFYGVKG	EKGDVGQPGP	NGIPSDTLHP	IIAPTGVTFH	PDQYKGEKGS
	······································	EGEPGIRGIS	LKGEEGIMGF PGL	PGLRGYPGLS (GEKGSPGQKG	SRGLDGYQGP	DGPRGPKGEA	GDPGPPGLPA
-		YSPHPSLAKG	ARGDPGFPGA	QGEPGSQGEP (GDPGLPGPPG	LSIGDGDQRR GLPGEMGPKG	GLPGEMGPKG	FIGDPGIPAL
		YGGPPGPDGK	RGPPGPPGLP	GPPGPDGFLF (GLKGAKGRAG	FPGLPGSPGA	RGPKGWKGDA	GECRCTEGDE
· · · · · · · · · · · · · · · · · · ·		AIKGLPGLPG	PKGFAGINGE	KGDKGDP (PGRKGDKGDP GQHGLPGFPG	LKGVPGNIGA	PGPKGAKGDS	RTITTKGERG
		QPGVPGVPGM	KGDDGSPGRD	GLDGFPGLPG	PPGDGIKGPP	GDPGYPGIPG	TKGTPGEMGP	PGLGLPGLKG
		QRGFPGDAGL	PGPPGFLGPP	GPAGTPGQID	CDTDVKRAVG	GDRQEAIQPG	CIAGPKGLPG	LPGPPGPTGA
		KGLRGIPGFA	GADGGPGPRG	LPGDAGREGF	PGPPGFIGPR	GSKGAVGLPG	PDGSPGPIGL	PGPDGPPGER
	·	GLPGEVLGAQ	PGPRGDAGVP	GOPGLKGLPG	DRGPPGFRGS	QGMPGMPGLK	GQPGLPGPSG	QPGLYGPPGL
***************************************		HGFPGAPGQE	GPLGLPGIPG	REGLPGDRGD	PGDTGAPGPV	GMKGLSGDRG	DAGFTGEQGH	PGSPGFKGID
		GMPGTPGLKG	DRGSPGMDGF	QGMPGLKGRP (GFPGSKGEAG	FFGIPGLKGL	FFGIPGLKGL AGEPGFKGSR	GDPGPPGPPP

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Seq.IDNO	Name	Proteinsequenz
		CTOCHIDATE GRADE
		VILPGMKDIK GEKGDEGPMG LKGYLGAKGI QGMPGIPGLS GIPGLPGRPG HIKGVKGDIG VPGIPGLPGF
		PGVAGPPGIT GFPGFIGSRG DKGAPGRAGL YGEIGATGDF GDIGDTINLP GRPGLKGERG TTGIPGLKGF
		FGEKGTEGDI GFPGITGVTG VQGPPGLKGQ TGFPGLTGPP GSQGELGRIG LPGGKGDDGW PGAPGLPGFP
		GLRGIRGLHG LPGTKGFPGS PGSDIHGDPG FPGPPGERGD PGEANTLPGP VGVPGQKGDQ GAPGERGPPG
		SPGLQGFPGI TPPSNISGAP GDKGAPGIFG LKGYRGPPGP PGSAALPGSK GDTGNPGAPG TPGTKGWAGD
		SGPQGRPGVF GLPGEKGPRG EQGFMGNTGP TGAVGDRGPK GPKGDPGFPG APGTVGAPGI AGIPQKIAIQ
		PGTVGPQGRR GPPGAPGEIG PQGPPGEPGF RGAPGKAGPQ GRGGVSAVPG FRGDEGPIGH QGPIGQEGAP
		GRPGSPGLPG MPGRSVSIGY LLVKHSQTDQ EPMCPVGMNK LWSGYSLLYF EGQEKAHNQD LGLAGSCLAR
		FSTMPFLYCN PGDVCYYASR NDKSYWLSTT APLPMMPVAE DEIKPYISRC SVCEAPAIAI AVHSQDVSIP
		HCPAGWRSLW IGYSFLMHTA AGDEGGGOSL VSPGSCLEDF RATPFIECNG GRGTCHYYAN KYSFWLTTIP
		EQSFQGSPSA DTLKAGLIRT HISRCQVCMK NL
		MERSALTSSA VISVAKVVRV ASGSAVVLPL ARIATVVIGG VVAMAAVPMV LSAMGFTAAG IASSSIAAKM
10	, 77 <u>4</u>	MSAAAIANGG GVASGSLVGT LQSLGATGLS GLTKFILGSI GSAIAAVIAR FY
,	מיולובמסווים ידים	MARGGRGRRL GLALGLILAL VLAPRVLRAK PTVRKERVVR PDSELGERPP EDNQSFQYDH EAFLGKEDSK
11	necicuiocaini	TPDOLTPDES KERLGKIVDR IDNDGDGFVT TEELKTWIKR VQKRYIFDNV AKVWKDYDRD KDDKISWEEY
		KOATYGYYLG NPAEFHDSSD HHTFKKMLPR DERRFKAADL NGDLTATREE FTAFLHPEEF EHMKEIVVLE
	2	TLEDIDKNGD GFVDQDEYIA DMFSHEENGP EPDWYLSERE QFNEFRDLNK DGKLDKDEIR HWILPQDYDH
		AQAEARHLVY ESDKNKDEKL TKEEILENWN MFVGSQATNY GEDLTKNHDE L
(Σ l dehvde	MATANGAVEN GQPDGKPPAL PRPIRNLEVK FTKIFINNEW HESKSGKKFA TCNPSTREQI CEVEEGDKPD
77	debydrogenase 6	VDKAVEAAQV AFQRGSPWRR LDALSRGRLL HQLADLVERD RATLAALETM DTGKPFLHAF FIDLEGCIRT

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LIRYFAGWADK IQGKTIPTDD LTALYLGSLI KEAGFPPGVV LGGKNPCIVC ADADLDLAVE TEQGPQIDQK QFDKILELIE KFKSIEBVIK RANSTDYGLT EYALAEYTEV KTVTIKLGDK MGAGSSTEQR SPEQPPEGSS QDELSLQEGD LNGQKGALNG ENRNIEQIPS SESNLEBLTQ AGDHQDPSLG AGEAASKESE ESPTSPVTSE TGSTFKKFFT ASEQAHPQEP AESAHEPRLS STVEERTEGQ KTEVEETAGS PPEGVVSEVE MLSSQERMKV KGESSASSPE EPEEITCLEK EDELDKVKSA TLSSTESTAS GGPKAMGGDH QKADBAGKDK SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKME	Proteinsequenz	
Gravin MG St		MILKDAROTTE AND TAGAT VERY MAIL KDAROTED
Lid	LRYFAGWADK IQGKTIPTDD NVVCFTRHEP IGVCGAITFW	NFFLEMLVWA LAFALCOGNI HVLNFALZIF
LIG TER Gravin MG Gravin PG BE BE GG GG GG GS	LIALYLGSLI KEAGFPPGVV NIVPGFGPTV GAAISSHPQI NKIAFTGSTE VGKLVKEAAS	NKIAFTGSTE VGKLVKEAAS RSNLKRVTLE
Gravin MG Gravin PE	LGGKNPCIVC ADADLDLAVE CAHQGVFFNQ GQCCTAASRV	FVEEQVYSEF VRRSVEYAKK RPVGDPFDVK
Gravin MG Gravin PG PI	OGPOIDOK OFDKILELIE SGKKEGAKLE	CGGSAMEDKG LFIKPTVFSE VTDNMRIAKE EIFGPVQPIL
Gravin MG OD PI EB	KFKSIEEVIK RANSTDYGLT AAVFTKNLDK ALKLASALES GTVWINCYNA LYAQAPFGGF	GTVWINCYNA LYAQAPFGGF KMSGNGRELG
Gravin Gravin Gravin Gravin GDELSLQEGD ENRNIEQIPS AGDHQDPSLG ESPTSPVTSE ASEQAHPQEP STVEERTEEQ PPEGVVSEVE KGESSASSPE EDELDKVKSA GGPKAMGGDH SKSKLEEKSE DVPAVVPLSE SWISASVTEP	EYALAEYTEV KTVTIKLGDK NP	
QDELSLQEGD ENRNIEQIPS AGDHQDPSLG ESPTSPVTSE ASEQAHPQEP STVEERTEEQ PPEGVVSEVE KGESSASSPE EDELDKVKSA GGPKAMGGDH SKSKLEEKSE DVPAVVPLSE SWISASVTEP	MGAGSSTEQR SPEQPPEGSS TPAEPEPSGG GPSAEAAPDT	TADPAIAASD PATKLLQKNG QLSTINGVAE
RNIEQIPS : DHQDPSLG PTSPVTSE PEQAHPQEP VEERTEEQ PEGVVSEVE BESSASSPE BESSASSPE	ELSLQEGD LNGQKGALNG QGALNSQEEE	EVIVTEVGQR DSEDVSERDS DKEMATKSAV VHDITDDGQE
AGDHQDPSLG AGEAASKESE ESPTSPVTSE TGSTFKKFFT ASEQAHPQEP AESAHEPRLS STVEERTEEQ KTEVEETAGS PPEGVVSEVE MLSSQERMKV KGESSASSPE BPEEITCLEK EDELDKVKSA TLSSTESTAE GGPKAMGGDH QKADEAGKDF SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKMI SWISASVTEP LEQVEAEAAI	ENRNIEQIPS SESNLEELTQ PTESQANDIG FKKVFKFVGF	KFTVKKDKTE KPDTVQLLTV KKDEGEGAAG
ESPTSPVTSE TGSTFKKFFT ASEQAHPQEP AESAHEPRLS STVEERTEEQ KTEVEETAGS PPEGVVSEVE MLSSQERMKV KGESSASSPE EPETTCLEK EDELDKVKSA TLSSTESTAS GGPKAMGGDH QKADEAGKDF SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKMI SWISASVTEP LEQVEAEAAI	AGDHQDPSLG AGBAASKESE PKQSTEKPEE TLKREQSHAE	ISPPAESGQA VEECKEEGEE KQEKEPSKSA
TEQAHPQEP TYEERTEEQ PEGVVSEVE BESSASSPE BELDKVKSA SPKAMGGDH CSKLEEKSE TPAVVPLSE	ESPTSPVTSE TGSTFKKFFT QGWAGWRKKT SFRKPKEDEV	EASEKKKEQE PEKVDTEEDG KAEVASEKLT
	ASEQAHPQEP AESAHEPRLS AEYEKVELPS EEQVSGSQGP	SEEKPAPLAT EVFDEKIEVH QEEVVAEVHV
	KTEVEETAGS VPAEELVGMD	AEPQEAEPAK ELVKLKETCV SGEDPTQGAD LSPDEKVLSK
	MLSSQERMKV QGSPLKKLFT	STGLKKLSGK KQKGKRGGGD EESGEHTQVP ADSPDSQEEQ
EDELDKVKSA TLSSTESTAS GGPKAMGGDH QKADEAGKDK SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKMF SWISASVTEP LEQVEAEAA	KGESSASSPE EPEEITCLEK GLAEVQQDGE AEEGATSDGE	KKREGVTPWA SFKKMVTPKK RVRRPSESDK
GGPKAMGGDH QKADEAGKDF SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKMF SWISASVTEP LEQVEAERAN	EDELDKVKSA TLSSTESTAS EMQEEMKGSV EEPKPEEPK	EEPKPEEPKR KVDTSVSWEA LICVGSSKKR ARRRSSSDEE
SKSKLEEKSE DSIAGSGVEF DVPAVVPLSE YDAVEREKMI SWISASVTEP LEQVEAEAN	GGPKAMGGDH QKADEAGKDK ETGTDGILAG SQEHDPGQGS	SSPEQAGSPT EGEGUSTWES FKRLVTPRKK
DVPAVVPLSE YDAVEREKMI SWISASVTEP LEQVEAEAAI	SKSKLEEKSE DSIAGSGVEH STPDTEPGKE ESWVSIKKFI	PGRRKKRPDG KQEQAPVEDA GPTGANEDDS
	DVPAVVPLSE YDAVEREKME AQQAQKGAEQ PEQKAATEVS	KELSESOVHM MAAAVADGTR AATIIEERSP
	SWISASVTEP LEQUEARAL LTEEVLEREV IAEEEPPTVT	EPLPENREAR GDTVVSEAEL TPEAVTAAET
AGPLGSEEGT EASAAEETTI	AGPLGSEEGT EASAAEETTE MVSAVSQLTD SPDTTEEATP VQEVEGGVPD	VQEVEGGVPD IEEQERRTQE VLQAVAEKVK

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Seq.IDNO	Name	Proteinsequenz
		CHRITYY YOO KE KOMMAGAS SASS
		EESQLPGTGG PEDVLQPVQR AEAERPEEQA EASGLKKEID VVLKVDAVEA KIEFILGGKV VGTIILDIE
		KAPQVTESIE SSELVTTCQA ETLAGVKSQE MVMEQAIPPD SVETPTDSET DGSTPVADFD APGTTQKDEI
		VEIHEENEVA SGTQSGGTEA EAVPAQKERP PAPSSFVFQE ETKEQSKMED TLEHTDKEVS VETVSILSKT
		EGTQEADQYA DEKTKDVPFF EGLEGSIDTG ITVSREKVTE VALKGEGTEE AECKKDDALE LQSHAKSPPS
		PVEREMYVQV EREKTEAEPT HVNEEKLEHE TAVTVSEEVS KQLLQTVNVP IIDGAKEVSS LEGSPPPCLG
- m. 1-11		QEEAVCTKIQ VQSSEASFTL TAAAEEEKVL GETANILETG ETLEPAGAHL VLEEKSSEKN EDFAAHPGED
		AVPTGPDCQA KSTPVIVSAT TKKGLSSDLE GEKTTSLKWK SDEVDEQVAC QEVKVSVAIE DLEPENGILE
		LETKSSKLVQ NIIQTAVDQF VRTEETATEM LISELQTQAH VIKADSQDAG QETEKEGEEP QASAQDETPI
		HSESDKAITP QAQEELQKQE RESAKSELTE S
7	Nidogen	MLASSSRIRA AWTRALLLPL LLAGPVGCLS RQELFPFGPG QGDLELEDGD DFVSPALELS GALRFYDRSD
ŗ,		IDAVYVTTNG IIATSEPPAK ESHPGLFPPT FGAVAPFLAD LDTTDGLGKV YYREDLSPSI TQRAAECVHR
		GPPEISFQPS SAVVVTWESV APYQGPSRDP DQKGKRNTFQ AVLASSDSSS YAIFLYPEDG LQFHTTFSKK
		ENNOVPAVVA FSQGSVGFLM KSNGAYNIFA NDRESIENLA KSSNSGQQGV WVFEIGSPAT TNGVVPADVI
		LGTEDGAEYD DEDEDYDLAT TRLGLEDVGT TPFSYKALRR GGADTYSVPS VLSPRRAATE RPLGPPTERT
		RSFQLAVETF HQQHPQVIDV DEVEETGVVF SYNTDSRQTC ANNRHQCSVH AECRDYATGF CCSCVAGYTG
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Seq.IDNO	Name	Proteinsequenz
		POSSITIED WELL VINCENCY THESETREYT VIRDERDIGAS PSRIYTYOWR OTITFOECUM DDSRPALPST
		YAFSNSIGPV REGSPDALQN PCYIGTHGCD
		IDECSEQPSV
		IPQRAQCIYT GGSSYTCSCL PGFSGDGQAC QDVDECQPSR CHPDAFCYNT PGSFTCQCKP GYQGDGFRCV
		PGEVEKTRCQ HEREHILGAA GATDPQRPIP PGLFVPECDA HGHYAPTQCH GSTGYCWCVD RDGREVEGTR
		TRPGMTPPCL STVAPPIHOG PAVPTAVIPL PPGTHLLFAQ TGKIERLPLE GNTMRKTEAK AFLHVPAKVI
	-	IGLAFDCVDK MVYWTDITEP SIGRASLHGG EPTTIIRQDL GSPEGIAVDH LGRNIFWTDS NLDRIEVAKL
		DGTQRRVLFE TDLVNPRGIV TDSVRGNLYW TDWNRDNPKI ETSYMDGTNR RILVQDDLGL PNGLHFDAFS
		SQLCWVDAGT NRAECLNPSQ PSRRKALEGL QYPFAVTSYG KNLYFTDWKM NSVVALDLAI SKETDAFQPH
		KQTRLYGITT ALSQCPQGHN YCSVNNGGCT HLCLATPGSR TCRCPDNTLG VDCIERK
1.5	Phospholipase C	MPSEKKISSA NDCISFMQAG CELKKVRPNS RIYNRFFTLD TDLQALRWEP SKKDLEKAKL DISAIKEIRL
n H	Ensilon	GKNTETFTNN GLADQICEDC AFSILHGENY ESLDLVANSA DVANIWVSGL RYLVSRSKQP LDFMEGNQNT
,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,		PRFMWLKTVF EAADVDGNGI MLEDTSVELI KQLNPTLKEA KIRLKFKEIQ KSKEKLTTRV TEEEFCEAFC
		ELCTRPEVYF LLVQISKNKE YLDANDLMLF LEAEQGVTHI TEDICLDIIR RYELSEEGRQ KGFLAIDGFT
		QYLLSSECDI FDPEQKKVAQ DMTQPLSHYY INASHNTYLI EDQFRGPADI NGYIRALKMG CRSVELDVSD
		GSDNEPILCN RNNMTTHVSF RSVIEVINKF AFVASEYPLI LCLGNHCSLP QQKVMAQQMK KVFGNKLYTE
		APLPSESYLP SPEKLKRMII VKGKKLPSDP DVLEGEVTDE DEEAQMSRRM SVDYNGEQKQ IRLCRELSDL
		VSICKSVQYR DFELSMKSQN YWEMCSFSET EASRIANEYP EDFVNYNKKF LSRIYPSAMR IDSSNLNPQD
		FWNCGCQIVA MNFQTPGPMM DLHTGWFLQN GGCGYVLRPS IMRDEVSYFS ANTKGILPGV SPLALHIKII
		SGONFPKPKG ACAKGDVIDP YVCIEIHGIP ADCSEORTKT VOONSDNPIF DETFEFOVNL PELAMIRFVV
		LDDDYIGDEF IGQYTIPFEC LOPGYRHVPL RSFVGDIMEH VTLFVHIAIT NRSGGGKAQK RSLSVRMGKK

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Seq.IDNO	Name	Proteinsequenz
		VREYTMLRNI GLKTIDDIFK IAVHPLREAI DMRENMONAI VSIKELCGLP PIASLKOCLL TLSSRLITSD
		NTPSVSLVMK DSFPYLEPLG AIPDVQKKML TAYDLMIQES RFLIEMADTV QEKIVQCQKA GMEFHEELHN
		LGAKEGLKGR KLNKATESFA WNITVLKGQG DLLKNAKNEA IENMKQIQLA CLSCGLSKAP SSSAEAKSKR
		SLEAIEEKES SEENGKL

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Proteinsequenz = Protein Sequence